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0570

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:	10/002,784	_
Source:	0185	_
Date Processed by STIC:	519/02	

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (http://www.uspto.gov/ebc/efs/downloads/documents.htm, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
- 3. Hand Carry directly to:
 U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name,
 Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
- U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1803, Crystal Plaza Two,
 2011 South Clark Place, Arlington, VA 22202
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 - Revised 01/29/2002



Does Not Comply Corrected Diskotte Needed

OIPE

DATE: 05/09/2002 RAW SEQUENCE LISTING TIME: 14:08:51 PATENT APPLICATION: US/10/002,784

Input Set : A:\Sequence.txt

Output Set: N:\CRF3\05092002\J002784.raw

gelete (3/33)9 <110> APPLICANT: Ulrich, Robert G. 11 <120> TITLE OF INVENTION: Bacterial Superantigen Vaccines 13 <130> FILE REFERENCE: 003/233/SAP 15 <140> CURRENT APPLICATION NUMBER: 10/002,784 C--> 17 <141> CURRENT FILING DATE: 2002-04-23 19 <150> PRIOR APPLICATION NUMBER: 08/882,431; 09/144,776 W--> 21 <151> PRIOR FILING DATE: 97-06-25; 98-09-01 23 <160> NUMBER OF SEQ ID NOS: 40 25 <170> SOFTWARE: Apple Macintosh Microsoft Word 6.0

ERRORED SEQUENCES

856 <210> SEQ ID NO: (17) 858 <211> LENGTH: 28 859 <212> TYPE: DNA 860 <213> ORGANISM: Arti/ficial sequence W--> 861 <220> FEATURE: 862 <223> OTHER INFORMATION: primer E--> 863 <400> SEQUENCE: (18) 865 ctcgcaagag gtacatatgc aacaagac 28 1121 <210> SEQ ID NO: 26 1123 <211> LENGTH: 220 1124 <212> TYPE: PRT 1125 <213> ORGANISM: Artificial sequence W--> 1126 <220> FEATURE: 1127 <223> OTHER INFORMATION: mutant streptococcal pyrogenic exotoxin-A W--> 1128 <400> SEQUENCE: 26 1130 Met Gln Gln Asp Pro Asp Pro Ser Gln Leu 1131 1132 His Arg Ser Ser Leu Val Lys Asn Leu Gln 1133 15 1134 Asn Ile Tyr Phe Leu Tyr Glu Gly Asp Pro 25 1136 Val Thr His Glu Asn Val Lys Ser Val Asp 35 1138 Gln Leu Arg Ser His Asp Leu Ile Tyr Asn 45 1140 Val Ser Gly Pro Asn Tyr Asp Lys Leu Lys 55

1143

1142 Thr Glu Leu Lys Asn Gln Glu Met Ala Thr

65

DATE: 05/09/2002

TIME: 14:08:51

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Input Set : A:\Sequence.txt
                                                         Output Set: N:\CRF3\05092002\J002784.raw
             1144 Leu Phe Lys Asp Lys Asn Ile Asp Ile Tyr
                                                                          75
             1146 Gly Val Glu Tyr Tyr His Leu Cys Tyr Leu
                                                                          85
             1148 Cys Glu Asn Ala Glu Arg Ser Ala Cys Ile
             1149
                                                                          95
             1150 Gly Gly Val Thr Asn Arg Glu Gly Asn His
                                                                                                                           (100
E--> 1151
                                                                       105
             1152 Leu Glu Ile Pro Lys Lys Ile Val Val Lys
                                                                       115
                                                                                                                              120
             1153
             1154 Val Ser Ile Asp Gly Ile Gln Ser Leu Ser
              1155
                                                                       125
             1156 Phe Asp Ile Glu Thr Asn Lys Lys Met Val
             1157
                                                                       135
             1158 Thr Ala Gln Glu Leu Asp Tyr Lys Val Arg
                                                                                                                              150
                                                                       145
             1160 Lys Tyr Leu Thr Asp Asn Lys Gln Leu Tyr
                                                                       155
              1161
              1162 Thr Asn Gly Pro Ser Lys Tyr Glu Thr Gly
                                                                                                                              170
              1163
                                                                       165
              1164 Tyr Ile Lys Phe Ile Pro Lys Asn Lys Glu
                                                                                                                              180
                                                                       175
                                                                                                                                                               The type of errors shown exist throughout
              1166 Ser Phe Trp Phe Asp Phe Phe Pro Glu Pro
                                                                                                                                                               the Sequence Listing. Please check subsequent
                                                                                                                              190
                                                                       185
                                                                                                                                                               sequences for similar errors.
              1168 Glu Phe Thr Gln Ser Lys Tyr Leu Met Ile
            220

220

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2208 Ever: 22087 is mondeten
              1169
                                                                       195
E--> 1330 <400> SEQUENCE: 32
              1336 Trp Tyr Asn Asp Leu Leu Val Asp Phe Asp
              1337
              1338 Ser Lys Asp Ile Val Asp Lys Tyr Lys Gly
              1339
              1340 Lys Lys Val Asp Leu Tyr Gly Ala Tyr Tyr
                                                                        45
              1342 Gly Tyr Gln Cys Ala Gly Gly Thr Pro Asn
              1343
                                                                        55
              1344 Lys Thr Ala Cys Met Tyr Gly Gly Val Thr
```

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/002,784

RAW SEQUENCE LISTING DATE: 05/09/2002 PATENT APPLICATION: US/10/002,784 TIME: 14:08:51

Input Set : A:\Sequence.txt

Output Set: N:\CRF3\05092002\J002784.raw

```
1345
     1346 Leu His Asp Asn Asn Arg Leu Thr Glu Glu
     1347
                           75
     1348 Lys Lys
     1350 <210> SEQ ID NO: 33
     1352 <211> LENGTH: 82
     1353 <212> TYPE: PRT
     1354 <213> ORGANISM: staphylococcal enterotoxin D
     1355 <223> OTHER INFORMATION: partial sequence as shown in Figure 3
E--> 1356 <400> SEQUENCE: 33
     1358 Thr Gly Asp Gln Phe Leu Glu Asn Thr Leu
     1360 Leu Tyr Lys Lys Phe Phe Thr Asp Leu Ile
     1361
     1362 Asn Phe Glu Asp Leu Leu Ile Asn Phe Asn
     1363
                           25
                                                30
     1364 Ser Lys Glu Met Ala Gln His Phe Lys Ser
     1365
                           35
     1366 Lys Asn Val Asp Val Tyr Pro Ile Arg Tyr
     1367
                           45
     1368 Ser Ile Asn Cys Tyr Gly Glu Ile Asp
                           55
     1370 Arg Thr Ala Cys Thr Tyr Gly Gly Val Thr
     1371 .
                          65
     1372 Pro His Glu Gly Asn Lys Leu Lys Glu Arg
     1373
     1374 Lys Lys
     1377 <210> SEQ ID NO: 34
     1379 <211> LENGTH: 82
     1380 <212> TYPE: PRT
     1381 <213> ORGANISM: staphylococcal enterotoxin E
     1382 <223> OTHER INFORMATION: partial sequence as shown in Figure 3
E--> 1383 <400> SEQUENCE: 34
     1385 Ser Asp Asp Gln Phe Leu Glu Asn Thr Leu
     1386
     1387 Leu Phe Lys Gly Phe Phe Thr Gly His Pro
     1388
     1389 Trp Tyr Asn Asp Leu Leu Val Asp Leu Gly
     1390
                          25
     1391 Ser Lys Asp Ala Thr Asn Lys Tyr Lys Gly
     1392
     1393 Lys Lys Val Asp Leu Tyr Gly Ala Tyr Tyr
     1394
                                                50
     1395 Gly Tyr Gln Cys Ala Gly Gly Thr Pro Asn
     1396
     1397 Lys Thr Ala Cys Met Tyr Gly Gly Val Thr
     1398
                          65
    1399 Leu His Asp Asn Asn Arg Leu Thr Glu Glu
     1400
                          75
```

DATE: 05/09/2002

TIME: 14:08:51

Input Set : A:\Sequence.txt Output Set: N:\CRF3\05092002\J002784.raw 1401 Lys Lys 1403 <210> SEQ ID NO: 35 1405 <211> LENGTH: 89 1406 <212> TYPE: PRT 1407 <213> ORGANISM: staphylococcal enterotoxin B 1408 <223> OTHER INFORMATION: partial sequence as shown in Figure 3 E--> 1409 <400> SEQUENCE: 35 1411 Ser Ile Asp Gln Phe Leu Tyr Phe Asp Leu 1412 1413 Ile Tyr Ser Ile Lys Asp Thr Lys Leu Gly 1415 Asn Tyr Asp Asn Val Arg Val Glu Phe Lys 1416 1417 Asn Lys Asp Leu Ala Asp Lys Tyr Lys Asp 1418 1419 Lys Tyr Val Asp Val Phe Gly Ala Asn Tyr 1420 50 45 1421 Tyr Gln Cys Tyr Phe Ser Lys Lys Thr Asn 1422 55 1423 Asp Ile Asn Ser His Gln Thr Asp Lys Arg 1424 65 1425 Lys Thr Cys Met Tyr Gly Gly Val Thr Glu 1426 75 1427 His Asn Gly Asn Gln Leu Asp Lys Tyr 1428 1430 <210> SEQ ID NO: 36 1432 <211> LENGTH: 89 1433 <212> TYPE: PRT 1434 <213> ORGANISM: staphylococcal enterotoxin C1 1435 <223> OTHER INFORMATION: partial sequence as shown in Figure 3 E--> 1436 <400> SEQUENCE: 36 1438 Ser Val Asp Lys Phe Leu Ala His Asp Leu 1440 Ile Tyr Asn Ile Ser Asp Lys Lys Leu Lys 1441 15 1442 Asn Tyr Asp Lys Val Lys Thr Glu Leu Leu 1443 25 1444 Asn Glu Gly Leu Ala Lys Lys Tyr Lys Asp 1445 35 40 1446 Glu Val Val Asp Val Tyr Gly Ser Asn Tyr 1447 1448 Tyr Val Asn Cys Tyr Phe Ser Ser Lys Asp 1449 60 1450 Asn Val Gly Lys Val Thr Gly Gly Lys Thr 1451 65 1452 Cys Met Tyr Gly Gly Ile Thr Lys His Glu 1454 Gly Asn His Phe Asp Asn Gly Asn Leu 1455

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/002,784

DATE: 05/09/2002

TIME: 14:08:51

Input Set : A:\Sequence.txt Output Set: N:\CRF3\05092002\J002784.raw 1457 <210> SEQ ID NO: 37 1459 <211> LENGTH: 89 1460 <212> TYPE: PRT 1461 <213> ORGANISM: staphylococcal enterotoxin C2 1462 <223> OTHER INFORMATION: partial sequence as shown in Figure 3 E--> 1463 <400> SEQUENCE: 37 1465 Ser Val Asp Lys Phe Leu Ala His Asp Leu 1467 Ile Tyr Asn Ile Ser Asp Lys Lys Leu Lys 1468 1469 Asn Tyr Asp Lys Val Lys Thr Glu Leu Leu 1470 1471 Asn Glu Asp Leu Ala Lys Lys Tyr Lys Asp 40 35 1472 1473 Glu Val Val Asp Val Tyr Gly Ser Asn Tyr 50 1474 45 1475 Tyr Val Asn Cys Tyr Phe Ser Ser Lys Asp 55 1476 1477 Asn Val Gly Lys Val Thr Gly Gly Lys Thr 65 1478 1479 Cys Met Tyr Gly Gly Ile Thr Lys His Glu 75 1481 Gly Asn His Phe Asp Asn Gly Asn Leu 1482 1484 <210> SEQ ID NO: 38 1486 <211> LENGTH: 89 1487 <212> TYPE: PRT 1488 <213> ORGANISM: staphylococcal enterotoxin C3 1489 <223> OTHER INFORMATION: partial sequence as shown in Figure 3 E--> 1490 <400> SEQUENCE: 38 1492 Ser Val Asp Lys Phe Leu Ala His Asp Leu 1493 1494 Ile Tyr Asn Ile Ser Asp Lys Lys Leu Lys 20 15 1495 1496 Asn Tyr Asp Lys Val Lys Thr Glu Leu Leu 25 1497 1498 Asn Glu Asp Leu Ala Lys Lys Tyr Lys Asp 35 1500 Glu Val Val Asp Val Tyr Gly Ser Asn Tyr 45 1501 1502 Tyr Val Asn Cys Tyr Phe Ser Ser Lys Asp 1503 1504 Asn Val Gly Lys Val Thr Gly Gly Lys Thr 1505 65 1506 Cys Met Tyr Gly Gly Ile Thr Lys His Glu 75 1508 Gly Asn His Phe Asp Asn Gly Asn Leu 1509 1511 <210> SEQ ID NO: 39

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/002,784

RAW SEQUENCE LISTING DATE: 05/09/2002 PATENT APPLICATION: US/10/002,784 TIME: 14:08:51

Input Set : A:\Sequence.txt

Output Set: N:\CRF3\05092002\J002784.raw

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1513 <211> LENGTH: 79
     1514 <212> TYPE: PRT
     1515 <213> ORGANISM: streptococcal pyrogenic enterotoxin a
     1516 <223> OTHER INFORMATION: partial sequence as shown in Figure 3
E--> 1517 <400> SEQUENCE: 39
     1519 Ser Val Asp Gln Leu Leu Ser His Asp Leu
     1520
     1521 Ile Tyr Asn Val Ser Gly Pro Asn Tyr Asp
     1523 Lys Leu Lys Thr Glu Leu Lys Asn Gln Glu
                          25
     1524
     1525 Met Ala Thr Leu Phe Lys Asp Lys Asn Val
     1526
     1527 Asp Ile Tyr Gly Val Glu Tyr Tyr His Leu
                                                50
     1528
                          45
     1529 Cys Tyr Leu Cys Glu Asn Ala Glu Arg Ser
     1530
     1531 Ala Cys Ile Tyr Gly Gly Val Thr Asn His
     1532
                          65
     1533 Glu Gly Asn His Leu Glu Ile Pro Lys
     1534
     1536 <210> SEQ ID NO: 40
     1538 <211> LENGTH: 73
     1539 <212> TYPE: PRT
     1540 <213> ORGANISM: toxin shock syndrome toxin-1
     1541 <223> OTHER INFORMATION: partial sequence as shown in Figure 3
E--> 1542 <400> SEQUENCE: 40
     1544 Val Leu Asp Asn Ser Leu Gly Ser Met Arg
     1545
     1546 Ile Lys Asn Thr Asp Gly Ser Ile Ser Leu
                                                20
                          15
     1548 Ile Ile Phe Pro Ser Pro Tyr Tyr Ser Pro
                           25
                                                30
     1549
     1550 Ala Phe Thr Lys Gly Glu Lys Val Asp Leu
     1551
                           35
     1552 Asn Thr Lys Arg Thr Lys Lys Ser Gln His
                           45
     1553
     1554 Thr Ser Glu Gly Thr Tyr Ile His Phe Gln
                           55
     1556 Ile Ser Gly Val Thr Asn Thr Glu Lys Leu
                                                70
     1557
     1558 Pro Thr Pro
```

VERIFICATION SUMMARYDATE: 05/09/2002PATENT APPLICATION: US/10/002,784TIME: 14:08:52

Input Set : A:\Sequence.txt

Output Set: N:\CRF3\05092002\J002784.raw

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L:3 M:259 W: Allowed number of lines exceeded, (1) GENERAL INFORMATION:
L:17 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:21 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:32 M:283 W: Missing Blank Line separator, <220> field identifier
L:34 M:283 W: Missing Blank Line separator, <400> field identifier
L:64 M:283 W: Missing Blank Line separator, <220> field identifier
L:66 M:283 W: Missing Blank Line separator, <400> field identifier
L:126 M:283 W: Missing Blank Line separator, <220> field identifier
L:128 M:283 W: Missing Blank Line separator, <400> field identifier
L:155 M:283 W: Missing Blank Line separator, <220> field identifier
L:214 M:283 W: Missing Blank Line separator, <220> field identifier
L:216 M:283 W: Missing Blank Line separator, <400> field identifier
L:267 M:283 W: Missing Blank Line separator, <220> field identifier
L:269 M:283 W: Missing Blank Line separator, <400> field identifier
L:331 M:283 W: Missing Blank Line separator, <220> field identifier
L:333 M:283 W: Missing Blank Line separator, <400> field identifier
L:384 M:283 W: Missing Blank Line separator, <220> field identifier
L:386 M:283 W: Missing Blank Line separator, <400> field identifier
L:448 M:283 W: Missing Blank Line separator, <220> field identifier
L:450 M:283 W: Missing Blank Line separator, <400> field identifier
L:494 M:283 W: Missing Blank Line separator, <220> field identifier
L:496 M:283 W: Missing Blank Line separator, <400> field identifier
L:552 M:283 W: Missing Blank Line separator, <220> field identifier
L:583 M:283 W: Missing Blank Line separator, <220> field identifier
L:585 M:283 W: Missing Blank Line separator, <400> field identifier
L:641 M:283 W: Missing Blank Line separator, <220> field identifier
L:643 M:283 W: Missing Blank Line separator, <400> field identifier
L:680 M:283 W: Missing Blank Line separator, <220> field identifier
L:682 M:283 W: Missing Blank Line separator, <400> field identifier
L:744 M:283 W: Missing Blank Line separator, <220> field identifier
L:746 M:283 W: Missing Blank Line separator, <400> field identifier
L:800 M:283 W: Missing Blank Line separator, <220> field identifier
L:802 M:283 W: Missing Blank Line separator, <400> field identifier
L:861 M:283 W: Missing Blank Line separator, <220> field identifier
L:863 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:17 differs:18
L:863 M:283 W: Missing Blank Line separator, <400> field identifier
L:872 M:283 W: Missing Blank Line separator, <220> field identifier L:874 M:283 W: Missing Blank Line separator, <400> field identifier
L:883 M:283 W: Missing Blank Line separator, <220> field identifier
L:885 M:283 W: Missing Blank Line separator, <400> field identifier
L:894 M:283 W: Missing Blank Line separator, <220> field identifier
L:896 M:283 W: Missing Blank Line separator, <400> field identifier
L:905 M:283 W: Missing Blank Line separator, <220> field identifier
L:907 M:283 W: Missing Blank Line separator, <400> field identifier
L:917 M:283 W: Missing Blank Line separator, <220> field identifier
L:919 M:283 W: Missing Blank Line separator, <400> field identifier
L:928 M:283 W: Missing Blank Line separator, <220> field identifier
L:930 M:283 W: Missing Blank Line separator, <400> field identifier
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VERIFICATION SUMMARY

DATE: 05/09/2002 TIME: 14:08:52

PATENT APPLICATION: US/10/002,784

Input Set : A:\Sequence.txt

Output Set: N:\CRF3\05092002\J002784.raw

L:976 M:283 W: Missing Blank Line separator, <220> field identifier L:978 M:283 W: Missing Blank Line separator, <400> field identifier L:1066 M:283 W: Missing Blank Line separator, <220> field identifier L:1068 M:283 W: Missing Blank Line separator, <400> field identifier L:1126 M:283 W: Missing Blank Line separator, <220> field identifier L:1128 M:283 W: Missing Blank Line separator, <400> field identifier L:1151 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:26 L:1330 M:200 E: Mandatory Header Field missing, <220> not found for SEQ ID#:32 L:1356 M:200 E: Mandatory Header Field missing, <220> not found for SEQ ID#:33 L:1383 M:200 E: Mandatory Header Field missing, <220> not found for SEQ ID#:34 L:1409 M:200 E: Mandatory Header Field missing, <220> not found for SEQ ID#:35 L:1436 M:200 E: Mandatory Header Field missing, <220> not found for SEQ ID#:36 L:1463 M:200 E: Mandatory Header Field missing, <220> not found for SEQ ID#:37 L:1490 M:200 E: Mandatory Header Field missing, <220> not found for SEQ ID#:38 L:1517 M:200 E: Mandatory Header Field missing, <220> not found for SEQ ID#:39 L:1542 M:200 E: Mandatory Header Field missing, <220> not found for SEQ ID#:40